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1: NM_017831. Homo sapiens hypo...[gi:8923428]

Related Sequences, Protein, Taxonomy, LinkOut

LOCUS NM_017831 1916 bp mRNA linear PRI 10-DEC-2001

DEFINITION Homo sapiens hypothetical protein FLJ20456 (FLJ20456), mRNA.

ACCESSION NM_017831

VERSION NM_017831.1 GI:8923428

2/15/00

KEYWORDS

SOURCE human.

ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (sites)

AUTHORS Watanabe,K., Kumagai,A., Itakura,S., Yamazaki,M., Tashiro,H.,
Ota,T., Suzuki,Y., Obayashi,M., Nishi,T., Shibahara,T., Tanaka,T.,
Nakamura,Y., Isogai,T. and Sugano,S.

TITLE NEDO human cDNA sequencing project

JOURNAL Unpublished (2000)

COMMENT PREDICTED REFSEQ: The mRNA record is supported by experimental
evidence; however, the coding sequence is predicted. The reference
sequence was derived from AK000463.1.

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1: AK000463. Homo sapiens cDNA...[gi:7020568]

Related Sequences, Protein, Taxonomy, LinkOut

LOCUS AK000463 1916 bp mRNA linear PRI 22-FEB-2000
DEFINITION Homo sapiens cDNA FLJ20456 fis, clone KAT05827.
ACCESSION AK000463
VERSION AK000463.1 GI:7020568
KEYWORDS oligo capping; fis (full insert sequence).
SOURCE Homo sapiens signet-ring cell carcinoma cell_line:KATO III cDNA to mRNA, clone_lib:KAT clone:KAT05827.
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (sites)
AUTHORS Watanabe,K., Kumagai,A., Itakura,S., Yamazaki,M., Tashiro,H., Ota,T., Suzuki,Y., Obayashi,M., Nishi,T., Shibahara,T., Tanaka,T., Nakamura,Y., Isogai,T. and Sugano,S.
TITLE NEDO human cDNA sequencing project
JOURNAL Unpublished (2000)
REFERENCE 2 (bases 1 to 1916)
AUTHORS Sugano,S., Suzuki,Y., Ota,T., Obayashi,M., Nishi,T., Isogai,T., Shibahara,T., Tanaka,T. and Nakamura,Y.
TITLE Direct Submission
JOURNAL Submitted (15-FEB-2000) Sumio Sugano, Institute of Medical Science, University of Tokyo, Deptment of Virology; Shirokane-dai, 4-6-1, Minato-ku, Tokyo 108-8639, Japan (E-mail:cdna1@ims.u-tokyo.ac.jp, Tel:81-3-5449-5286, Fax:81-3-5449-5416)
COMMENT NEDO human cDNA sequencing project supported by Ministry of International Trade and Industry of Japan; cDNA full insert sequencing: Research Association for Biotechnology; cDNA library construction, 5' & 3'-end one pass sequencing: Departent of Virology and Human Genome Center, Institute of Medical Science, University of Tokyo (partly supported by Science and Technology Agency).
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1: XM_008732. Homo sapiens hypo...[gi:18588856]

Related Sequences. Protein, Taxonomy, LinkOut

LOCUS XM_008732 1916 bp mRNA linear PRI 07-FEB-2002
DEFINITION Homo sapiens hypothetical protein FLJ20456 (FLJ20456), mRNA.
ACCESSION XM_008732
VERSION XM_008732.4 GI:18588856
KEYWORDS
SOURCE human.
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 1916)
AUTHORS NCBI Annotation Project.
TITLE Direct Submission
JOURNAL Submitted (06-FEB-2002) National Center for Biotechnology
Information, NIH, Bethesda, MD 20894, USA
COMMENT GENOME ANNOTATION REFSEQ: This model reference sequence was
predicted from NCBI contig NT_010966 by automated computational
analysis using gene prediction method: BLAST. ~Also see:-
Documentation of NCBI's Annotation Process - Evidence Viewer -
alignments supporting this model.
On Feb 7, 2002 this sequence version replaced gi:17456245.
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